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,	VEGF-B167		v		_	_	s -		-	-		· , -		R	- 1																R C		160
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	VEGF-C VEGF-D		Ö																						ME						CV		273

Fig. 2A

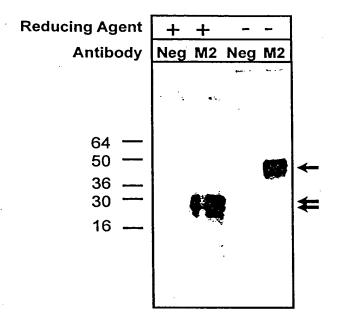


Fig. 2B

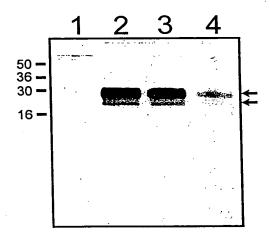
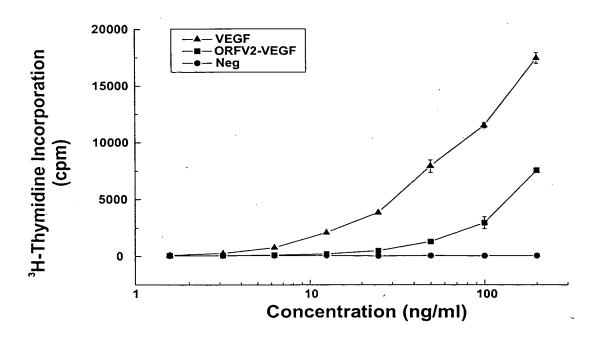


Fig. 3





9

Fig. 4A

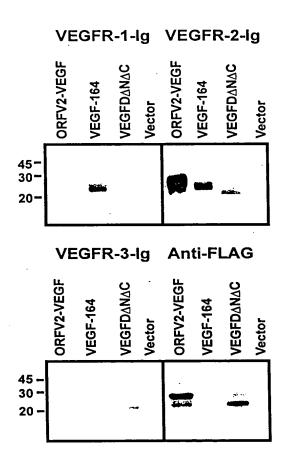


Fig. 4B

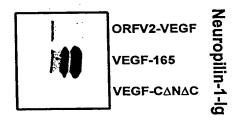


Fig. 5

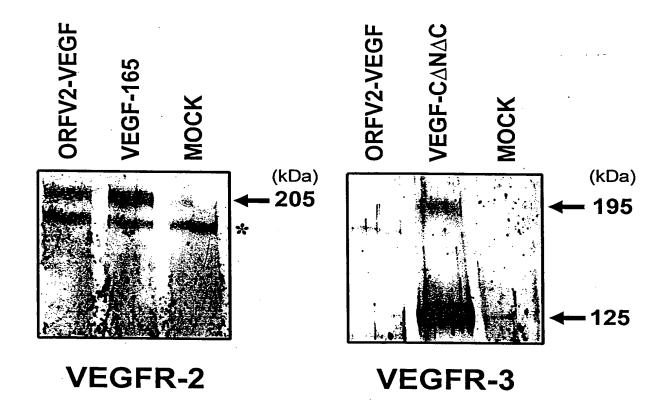


Fig. 6

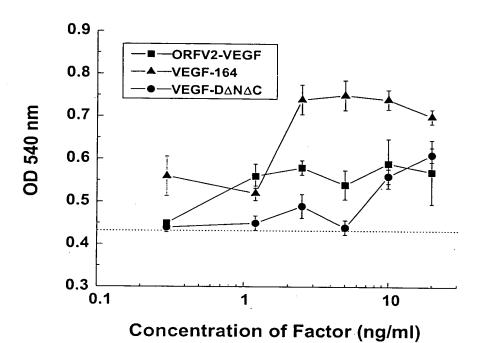


Fig. 7A

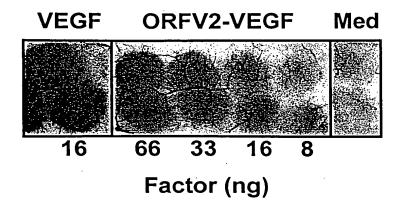


Fig. 7B

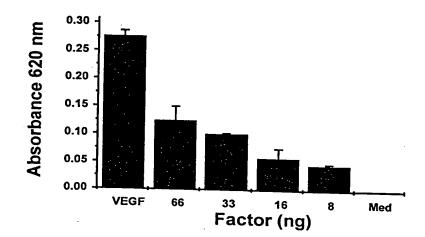


Fig. 8

Fig. 9

MKLLVGILVAVCLHQYLLNADSNTKGWSEVLKGSECKPRPIVVPVSETHPELTSQRFNPPCVTLM RCGGCCNDESLECVPTEEVNVTMELLGASGSGSNGMQRLSFVEHKKCDCRPRFTTTPPTTTRPPR RRR

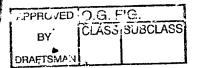


Fig. 10

atgaagttgc tcgtcggcat actggtagcc gtgtgcttgc accagtatct gctgaacgcg 60 gacagcacga aaacatggtc cgaggtgttt gaaagcagta agtgcaagcc aaggccaacg 120 gtcgttcccg taggcgaggc gcacccagag ctaacttctc agcggttcaa cccgcagtgt 180 gtcacagtga tgcgatgcgg cgggtgctgc aacgacgaga gcttggaatg cgtccccacg 240 gaagaggcaa acgtgacgat gcaactcatg ggggcgtcgg tctccggtgg taacgggatg 300 caacatttga tattcgtaga gcataagaaa tgcgattgta aaccacgact cacaaccacg 360 ccaccgacga ccacaaggcc gcccagaaga cgccgctag

Fig. 11

Met Lys Leu Val Gly Ile Leu Val Ala Val Cys Leu His Gln Tyr
1 5 10 15

Leu Leu Asn Ala Asp Ser Thr Lys Thr Trp Ser Glu Val Phe Glu Ser 20 25 30

Ser Lys Cys Lys Pro Arg Pro Thr Val Val Pro Val Gly Glu Ala His $35 \hspace{1cm} 40 \hspace{1cm} 45$

Pro Glu Leu Thr Ser Gln Arg Phe Asn Pro Gln Cys Val Thr Val Met 50 55 60

Arg Cys Gly Gly Cys Cys Asn Asp Glu Ser Leu Glu Cys Val Pro Thr
65 70 75 80

Glu Glu Ala Asn Val Thr Met Gln Leu Met Gly Ala Ser Val Ser Gly 85 90 95

Gly Asn Gly Met Gln His Leu Ile Phe Val Glu His Lys Lys Cys Asp 100 105 110

Cys Lys Pro Arg Leu Thr Thr Thr Pro Pro Thr Thr Thr Arg Pro Pro 115 120 125

Arg Arg Arg Arg 130